```
        RAW SEQUENCE LISTING                  DATE: 01/04/2001
        PATENT APPLICATION:  US/09/673,198    TIME: 11:31:43

        Input Set : A:\Pto.amc
        Output Set: N:\CRF3\01042001\I673198.raw

     2 <110> APPLICANT: MIYAKE, Koichiro; HASHIMOTO, Shinichi; MOTOYAMA Hiroaki;
     3         OZAKI, Akio; SETO, Haruo; KUZAYAMA, Tomohisa; TAKAHASHI, Shunji
     5 <120> TITLE OF INVENTION: A process for producing isoprenoid compounds by
     6         microorganisms and a method for screening compounds with
     7         antibiotic or weeding activity
W--> 9 <130> FILE REFERENCE:
C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/673,198
C--> 12 <141> CURRENT FILING DATE: 2000-10-12
    14 <150> PRIOR APPLICATION NUMBER: JP98/103101
    15 <151> PRIOR FILING DATE: 1998-04-14
    17 <150> PRIOR APPLICATION NUMBER: JP98/221910
    18 <151> PRIOR FILING DATE: 1998-08-05
    20 <150> PRIOR APPLICATION NUMBER: JP99/035739
    21 <151> PRIOR FILING DATE: 1999-02-15
    23 <160> NUMBER OF SEQ ID NOS: 34
    25 <170> SOFTWARE: PatentIn Ver. 2.0
    27 <210> SEQ ID NO: 1
    28 <211> LENGTH: 620
    29 <212> TYPE: PRT
    30 <213> ORGANISM: Escherichia coli
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    36 Thr Gln Glu Leu Arg Leu Leu Pro Lys Glu Ser Leu Pro Lys Leu Cys
    37             20              25              30
    39 Asp Glu Leu Arg Arg Tyr Leu Leu Asp Ser Val Ser Arg Ser Ser Gly
    40         35              40              45
    42 His Phe Ala Ser Gly Leu Gly Thr Val Glu Leu Thr Val Ala Leu His
    43         50              55              60
    45 Tyr Val Tyr Asn Thr Pro Phe Asp Gln Leu Ile Trp Asp Val Gly His
    46  65              70              75              80
    48 Gln Ala Tyr Pro His Lys Ile Leu Thr Gly Arg Arg Asp Lys Ile Gly
    49             85              90              95
    51 Thr Ile Arg Gln Lys Gly Gly Leu His Pro Phe Pro Trp Arg Gly Glu
    52             100             105             110
    54 Ser Glu Tyr Asp Val Leu Ser Val Gly His Ser Ser Thr Ser Ile Ser
    55             115             120             125
    57 Ala Gly Ile Gly Ile Ala Val Ala Ala Glu Lys Glu Gly Lys Asn Arg
    58             130             135             140
    60 Arg Thr Val Cys Val Ile Gly Asp Gly Ala Ile Thr Ala Gly Met Ala
    61 145             150             155             160
    63 Phe Glu Ala Met Asn His Ala Gly Asp Ile Arg Pro Asp Met Leu Val
    64             165             170             175
    66 Ile Leu Asn Asp Asn Glu Met Ser Ile Ser Glu Asn Val Gly Ala Leu
    67             180             185             190
    69 Asn Asn His Leu Ala Gln Leu Leu Ser Gly Lys Leu Tyr Ser Ser Leu
    70             195             200             205
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RAW SEQUENCE LISTING   DATE: 01/04/2001
PATENT APPLICATION: US/09/673,198   TIME: 11:31:43

Input Set : A:\Pto.amc
Output Set: N:\CRF3\01042001\I673198.raw

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 72 Arg Glu Gly Gly Lys Lys Val Phe Ser Gly Val Pro Pro Ile Lys Glu
 73     210                 215                 220
 75 Leu Leu Lys Arg Thr Glu Glu His Ile Lys Gly Met Val Val Pro Gly
 76 225                 230                 235                 240
 78 Thr Leu Phe Glu Glu Leu Gly Phe Asn Tyr Ile Gly Pro Val Asp Gly
 79                 245                 250                 255
 81 His Asp Val Leu Gly Leu Ile Thr Thr Leu Lys Asn Met Arg Asp Leu
 82                 260                 265                 270
 84 Lys Gly Pro Gln Phe Leu His Ile Met Thr Lys Lys Gly Arg Gly Tyr
 85             275                 280                 285
 87 Glu Pro Ala Glu Ly  Asp Pro Ile Thr Phe His Ala Val Pro Lys Phe
 88     290                 295                 300
 90 Asp Pro Ser Ser Gly Cys Leu Pro Lys Ser Ser Gly Gly Leu Pro Ser
 91 305                 310                 315                 320
 93 Tyr Ser Lys Ile Phe Gly Asp Trp Leu Cys Glu Thr Ala Ala Lys Asp
 94                 325                 330                 335
 96 Asn Lys Leu Met Ala Ile Thr Pro Ala Met Arg Glu Gly Ser Gly Met
 97                 340                 345                 350
 99 Val Glu Phe Ser Arg Lys Phe Pro Asp Arg Tyr Phe Asp Val Ala Ile
100                 355                 360                 365
102 Ala Glu Gln His Ala Val Thr Phe Ala Ala Gly Leu Ala Ile Gly Gly
103     370                 375                 380
105 Tyr Lys Pro Ile Val Ala Ile Tyr Ser Thr Phe Leu Gln Arg Ala Tyr
106 385                 390                 395                 400
108 Asp Gln Val Leu His Asp Val Ala Ile Gln Lys Leu Pro Val Leu Phe
109                 405                 410                 415
111 Ala Ile Asp Arg Ala Gly Ile Val Gly Ala Asp Gly Gln Thr His Gln
112             420                 425                 430
114 Gly Ala Phe Asp Leu Ser Tyr Leu Arg Cys Ile Pro Glu Met Val Ile
115             435                 440                 445
117 Met Thr Pro Ser Asp Glu Asn Glu Cys Arg Gln Met Leu Tyr Thr Gly
118     450                 455                 460
120 Tyr His Tyr Asn Asp Gly Pro Ser Ala Val Arg Tyr Pro Arg Gly Asn
121 465                 470                 475                 480
123 Ala Val Gly Val Glu Leu Thr Pro Leu Glu Lys Leu Pro Ile Gly Lys
124                 485                 490                 495
126 Gly Ile Val Lys Arg Arg Gly Glu Lys Leu Ala Ile Leu Asn Phe Gly
127             500                 505                 510
129 Thr Leu Met Pro Glu Ala Ala Lys Val Ala Glu Ser Leu Asn Ala Thr
130             515                 520                 525
132 Leu Val Asp Met Arg Phe Val Lys Pro Leu Asp Glu Ala Leu Ile Leu
133     530                 535                 540
135 Glu Met Ala Ala Ser His Glu Ala Leu Val Thr Val Glu Glu Asn Ala
136 545                 550                 555                 560
138 Ile Met Gly Gly Ala Gly Ser Gly Val Asn Glu Val Leu Met Ala His
139                 565                 570                 575
141 Arg Lys Pro Val Pro Val Leu Asn Ile Gly Leu Pro Asp Phe Phe Ile
142     580                 585                 590
144 Pro Gln Gly Thr Gln Glu Glu Met Arg Ala Glu Leu Gly Leu Asp Ala
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/673,198

Input Set : A:\Pto.amc
Output Set: N:\CRF3\01042001\I673198.raw

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154 <213> ORGANISM: Escherichia coli
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160 Ala Leu Ser Arg Phe Ile Ala Pro Leu Pro Phe Gln Asn Thr Pro Val
161                20                  25                 30
163 Val Glu Thr Met Gln Tyr Gly Ala Leu Leu Gly Gly Lys Arg Leu Arg
164            35                  40                  45
166 Pro Phe Leu Val Tyr Ala Thr Gly His Met Phe Gly Val Ser Thr Asn
167      50                  55                  60
169 Thr Leu Asp Ala Pro Ala Ala Ala Val Glu Cys Ile His Ala Tyr Ser
170   65                  70 .                 75                 80
172 Leu Ile His Asp Asp Leu Pro Ala Met Asp Asp Asp Leu Arg Arg
173            85                  90                  95
175 Gly Leu Pro Thr Cys His Val Lys Phe Gly Glu Ala Asn Ala Ile Leu
176            100                 105                 110
178 Ala Gly Asp Ala Leu Gln Thr Leu Ala Phe Ser Ile Leu Ser Asp Ala
179            115                 120                 125
181 Asp Met Pro Glu Val Ser Asp Arg Asp Arg Ile Ser Met Ile Ser Glu
182      130                 135                 140
184 Leu Ala Ser Ala Ser Gly Ile Ala Gly Met Cys Gly Gly Gln Ala Leu
185 145                 150                 155                 160
187 Asp Leu Asp Ala Glu Gly Lys His Val Pro Leu Asp Ala Leu Glu Arg
188            165                 170                 175
190 Ile His Arg His Lys Thr Gly Ala Leu Ile Arg Ala Ala Val Arg Leu
191            180                 185                 190
193 Gly Ala Leu Ser Ala Gly Asp Lys Gly Arg Arg Ala Leu Pro Val Leu
194            195                 200                 205
196 Asp Lys Tyr Ala Glu Ser Ile Gly Leu Ala Phe Gln Val Gln Asp Asp
197      210                 215                 220
199 Ile Leu Asp Val Val Gly Asp Thr Ala Thr Leu Gly Lys Arg Gln Gly
200 225                 230                 235                 240
202 Ala Asp Gln Gln Leu Gly Lys Ser Thr Tyr Pro Ala Leu Leu Gly Leu
203            245.                250                 255
205 Glu Gln Ala Arg Lys Lys Ala Arg Asp Leu Ile Asp Asp Ala Arg Gln
206            260                 265                 270
208 Ser Leu Lys Gln Leu Ala Glu Gln Ser Leu Asp Thr Ser Ala Leu Glu
209         275                 280                 285
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216 <211> LENGTH: 80
217 <212> TYPE: PRT
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225             20                   25                  30
227 Glu Glu Ala Leu Asn Glu Phe Glu Arg Gly Val Gln Leu Ala Arg Gln
228             35                   40                  45
230 Gly Gln Ala Lys Leu Gln Gln Ala Glu Gln Arg Val Gln Ile Leu Leu
231         50                  55                  60
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234  65                  70                  75                  80
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238 <211> LENGTH: 348
239 <212> TYPE: PRT
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247             20                   25                  30
249 Thr Asp Leu Arg Val Ser Arg Leu Cys Leu Gly Cys Met Thr Phe Gly
250             35                   40                  45
252 Glu Pro Asp Arg Gly Asn His Ala Trp Thr Leu Pro Glu Glu Ser Ser
253         50                  55                  60
255 Arg Pro Ile Ile Lys Arg Ala Leu Glu Gly Gly Ile Asn Phe Phe Asp
256  65                  70                  75                  80
258 Thr Ala Asn Ser Tyr Ser Asp Gly Ser Ser Glu Glu Ile Val Gly Arg
259                 85                  90                  95
261 Ala Leu Arg Asp Phe Ala Arg Arg Glu Asp Val Val Val Ala Thr Lys
262                 100                 105                 110
264 Val Phe His Arg Val Gly Asp Leu Pro Glu Gly Leu Ser Arg Ala Gln
265             115                 120                 125
267 Ile Leu Arg Ser Ile Asp Asp Ser Leu Arg Arg Leu Gly Met Asp Tyr
268     130                 135                 140
270 Val Asp Ile Leu Gln Ile His Arg Trp Asp Tyr Asn Thr Pro Ile Glu
271 145                 150                 155                 160
273 Glu Thr Leu Glu Ala Leu Asn Asp Val Val Lys Ala Gly Lys Ala Arg
274             165                 170                 175
276 Tyr Ile Gly Ala Ser Ser Met His Ala Ser Gln Phe Ala Gln Ala Leu
277             180                 185                 190
279 Glu Leu Gln Lys Gln His Gly Trp Ala Gln Phe Val Ser Met Gln Asp
280             195                 200                 205
282 His Tyr Asn Leu Ile Tyr Arg Glu Glu Glu Arg Glu Met Leu Pro Leu
283     210                 215                 220
285 Cys Tyr Gln Glu Gly Val Ala Val Ile Pro Trp Ser Pro Leu Ala Arg
286 225                 230                 235                 240
288 Gly Arg Leu Thr Arg Pro Trp Gly Glu Thr Thr Ala Arg Leu Val Ser
289             245                 250                 255

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291 Asp Glu Val Gly Lys Asn Leu Tyr Lys Glu Ser Asp Glu Asn Asp Ala
292             260             265             270
294 Gln Ile Ala Glu Arg Leu Thr Gly Val Ser Glu Glu Leu Gly Ala Thr
295         275             280             285
297 Arg Ala Gln Val Ala Leu Ala Trp Leu Leu Ser Lys Pro Gly Ile Ala
298     290             295             300
300 Ala Pro Ile Ile Gly Thr Ser Arg Glu Glu Gln Leu Asp Glu Leu Leu
301 305             310             315             320
303 Asn Ala Val Asp Ile Thr Leu Lys Pro Glu Gln Ile Ala Glu Leu Glu
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311 <211> LENGTH: 398
312 <212> TYPE: PRT
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319 Thr Leu Asp Val Val Arg His Asn Pro Glu His Phe Arg Val Val Ala
320             20              25              30
322 Leu Val Ala Gly Lys Asn Val Thr Arg Met Val Glu Gln Cys Leu Glu
323         35              40              45
325 Phe Ser Pro Arg Tyr Ala Val Met Asp Asp Glu Ala Ser Ala Lys Leu
326     50              55              60
328 Leu Lys Thr Met Leu Gln Gln Gly Ser Arg Thr Glu Val Leu Ser
329 65              70              75              80
331 Gly Gln Gln Ala Ala Cys Asp Met Ala Ala Leu Glu Asp Val Asp Gln
332             85              90              95
334 Val Met Ala Ala Ile Val Gly Ala Ala Gly Leu Leu Pro Thr Leu Ala
335             100             105             110
337 Ala Ile Arg Ala Gly Lys Thr Ile Leu Leu Ala Asn Lys Glu Ser Leu
338         115             120             125
340 Val Thr Cys Gly Arg Leu Phe Met Asp Ala Val Lys Gln Ser Lys Ala
341     130             135             140
343 Gln Leu Leu Pro Val Asp Ser Glu His Asn Ala Ile Phe Gln Ser Leu
344 145             150             155             160
346 Pro Gln Pro Ile Gln His Asn Leu Gly Tyr Ala Asp Leu Glu Gln Asn
347             165             170             175
349 Gly Val Val Ser Ile Leu Leu Thr Gly Ser Gly Gly Pro Phe Arg Glu
350             180             185             190
352 Thr Pro Leu Arg Asp Leu Ala Thr Met Thr Pro Asp Gln Ala Cys Arg
353         195             200             205
355 His Pro Asn Trp Ser Met Gly Arg Lys Ile Ser Val Asp Ser Ala Thr
356     210             215             220
358 Met Met Asn Lys Gly Leu Glu Tyr Ile Glu Ala Arg Trp Leu Phe Asn
359 225             230             235             240
361 Ala Ser Ala Ser Glu Met Glu Val Leu Ile His Pro Gln Ser Val Ile
362             245             250             255
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VERIFICATION SUMMARY   DATE: 01/04/2001
PATENT APPLICATION: US/09/673,198   TIME: 11:31:44

Input Set : A:\Pto.amc
Output Set: N:\CRF3\01042001\I673198.raw

L:9 M:201 W: Mandatory field data missing, FILE REFERENCE
L:11 M:270 C: Current Application Number differs, Replaced Current Application Number
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:1054 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 11
L:1055 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1060 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1065 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1070 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1075 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
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L:1270 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/673,198

DATE: 01/04/2001
TIME: 11:31:44

Input Set : A:\Pto.amc
Output Set: N:\CRF3\01042001\I673198.raw

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L:1275 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1280 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
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L:1295 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1300 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
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Serial Number: ●/673,198   **ENTERED**   CRF Processing Date: 12/26/~~2~~00
Edited by:
Verified by: AL   (STIC

☐ Changed a file from non-ASCII to ASCII

☑ Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☐ Edited a format error in the Current Application Data section, specifically:

_____

☑ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.

☐ Added the mandatory heading and subheadings for "Current Application Data".

☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____

☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:

_____

☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

_____

☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐ Inserted colons after headings/subheadings. Headings edited included:

_____

☐ Deleted extra, invalid, headings used by an applicant, specifically:

_____

☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of t
☐ page numbers throughout text; ☐ other invalid text, such as_____

☐ Inserted mandatory headings, specifically: _____

☐ Corrected an obvious error in the response, specifically:

_____

☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

☐ Corrected an error in the Number of Sequences field, specifically: _____

☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (errc
due to a PatentIn bug). Sequences corrected: _____

☐ Other:

_____

_____

**Examiner: The above corrections must be communicated to the applicant in the first Office
Action. DO NOT send a copy of this form.**   3/1/95